## NOV 0 6 2000 30 HO

## SEQUENCE LISTING

<110> A Para Bio ciences Corporation

Zucker Maarles Mendlein, John Sun, Yumei Tsunoda, Susan Sierralta, Jimena

<120> Compositions And Methods For Identifying Modulators and Transducisomes

<130> AURO1210-1

<140> 09/462,517

<141> 2000-05-18

<160> 16

<170> PatentIn version 3.0

<210> 1

<211> 674

<212> PRT

<213> Drosophila melanogaster

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Val Arg Gly Glu Val Lys Asp Ser Pro Asn Thr Lys Thr Thr Gly Ile 35 40 45

Phe Ile Lys Gly Ile Val Pro Asp Ser Pro Ala His Leu Cys Gly Arg
50 55 60

Leu Lys Val Gly Asp Arg Ile Leu Ser Leu Asn Gly Lys Asp Val Arg 65 70 75 80

Asn Ser Thr Glu Gln Ala Val Ile Asp Leu Ile Lys Glu Ala Asp Phe 85 90 95

Lys Ile Glu Leu Glu Ile Gln Thr Phe Asp Lys Ser Asp Glu Gln Gln 100 105 110

Ala Lys Ser Asp Pro Arg Ser Asn Gly Tyr Met Gln Ala Lys Asn Lys
115 120 125

Phe Asn Gln Glu Gln Thr Thr Asn Asn Asn Ala Ser Gly Gln Gly
130 135 140

Met Gly Gln Gly Gln Gly Gln Gly Met Ala Gly Met Asn Arg 145 150 155 160

Gln Gln Ser Met Gln Lys Arg Asn Thr Thr Phe Thr Ala Ser Met Arg 165 170 175

Gln Lys His Ser Asn Tyr Ala Asp Glu Asp Asp Glu Asp Thr Arg Asp 180 185 190

Met Thr Gly Arg Ile Arg Thr Glu Ala Gly Tyr Glu Ile Asp Arg Ala 195 200 205

Ser Ala Gly Asn Cys Lys Leu Asn Lys Gln Glu Lys Asp Arg Asp Lys Glu Gln Glu Asp Glu Phe Gly Tyr Thr Met Ala Lys Ile Asn Lys Arg Tyr Asn Met Met Lys Asp Leu Arg Arg Ile Glu Val Gln Arg Asp Ala 250 Ser Lys Pro Leu Gly Leu Ala Leu Ala Gly His Lys Asp Arg Gln Lys Met Ala Cys Phe Val Ala Gly Val Asp Pro Asn Gly Ala Leu Gly Ser Val Asp Ile Lys Pro Gly Asp Glu Ile Val Glu Val Asn Gly Asn Val 295 Leu Lys Asn Arg Cys His Leu Asn Ala Ser Ala Val Phe Lys Asn Val Asp Gly Asp Lys Leu Val Met Ile Thr Ser Arg Arg Lys Pro Asn Asp 330 Glu Gly Met Cys Val Lys Pro Ile Lys Lys Phe Pro Thr Ala Ser Asp Glu Thr Lys Phe Ile Phe Asp Gln Phe Pro Lys Ala Arg Thr Val Gln 360 Val Arg Lys Glu Gly Phe Leu Gly Ile Met Val Ile Tyr Gly Lys His Ala Glu Val Gly Ser Gly Ile Phe Ile Ser Asp Leu Arg Glu Gly Ser 390 Asn Ala Glu Leu Ala Gly Val Lys Val Gly Asp Met Leu Leu Ala Val 410 Asn Gln Asp Val Thr Leu Glu Ser Asn Tyr Asp Asp Ala Thr Gly Leu 425 Leu Lys Arg Ala Glu Gly Val Val Thr Met Ile Leu Leu Thr Leu Lys 440 Ser Glu Glu Ala Ile Lys Ala Glu Lys Ala Ala Glu Glu Lys Lys Glu Glu Ala Lys Lys Glu Glu Glu Lys Pro Gln Glu Pro Ala Thr Ala Glu Ile Lys Pro Asn Lys Lys Ile Leu Ile Glu Leu Lys Val Glu Lys 490 Lys Pro Met Gly Cys His Arg Leu Arg Arg Gln Lys Gln Pro Cys His Asp Trp Leu Cys Asn His Pro Arg Leu Ser Gly Gly Gln Val Ala Ala 520 Asp Lys Arg Leu Lys Ile Phe Asp His Ile Cys Asp Ile Asn Gly Thr Pro Ile His Val Gly Ser Met Thr Thr Leu Lys Val His Gln Leu Phe 545 550 555 560

His Thr Tyr Glu Lys Ala Val Thr Leu Thr Val Phe Arg Ala Asp 565 570 575

Pro Pro Glu Leu Glu Lys Phe Asn Val Asp Leu Met Lys Lys Ala Gly 580 585 590

Lys Glu Leu Gly Leu Ser Leu Ser Pro Asn Glu Ile Gly Cys Thr Ile
595 600 605

Ala Asp Leu Ile Gln Gly Gln Tyr Pro Glu Ile Asp Ser Lys Leu Gln 610 620

Arg Gly Asp Ile Ile Thr Lys Phe Asn Gly Asp Ala Leu Glu Gly Leu 625 630 635

Pro Phe Gln Val Cys Tyr Ala Leu Phe Lys Gly Ala Asn Gly Lys Val 645 . 650 655

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Phe Ser Ile Ala Gly Gly Thr Asp Asn Pro His Ile Gly Asp Asp Pro 20 25 30

Ser Ile Phe Ile Thr Lys Ile Ile Pro Gly Gly Ala Ala Ala Gln Asp 35 40 45

Gly Arg Leu Arg Val Asn Asp Ser Ile Leu Phe Val Asn Glu Val Asp 50 60

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Gly Ser Ile Val Arg Leu Tyr Val Met Arg Arg Lys Pro

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Ser Ile Tyr Val Thr Lys Ile Ile Glu Gly Gly Ala Ala His Lys Asp
Gly Arg Leu Gln Ile Gly Asp Lys Ile Leu Ala Val Asn Ser Val Gly
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Tyr Asp Val Val Tyr Leu Lys Val Ala Lys Pro Ser Asn
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Asp Gln Ile Leu Ser Val Asn Gly Val Asp Leu Arg Asn Ala Ser His
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Met Val Ile Tyr Gly Lys His Ala Glu Val Gly Ser Gly Ile Phe Ile

Ser Asp Leu Arg Glu Gly Ser Asn Ala Glu Leu Ala Gly Val Lys Val

35 40 45

Gly Asp Met Leu Leu Ala Val Asn Gln Asp Val Thr Leu Glu Ser Asn 50 60

Tyr Asp Asp Ala Thr Gly Leu Leu Lys Arg Ala Glu Gly Val Val Thr 65 70 75 80

Met Ile Leu Leu Thr Leu Lys Ser

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Gly Ile Cys Ile Val Arg Gly Glu Val Lys Asp Ser Pro Asn Thr Lys
20 25 30

Thr Thr Gly Ile Phe Ile Lys Gly Ile Val Pro Asp Ser Pro Ala His

Leu Cys Gly Arg Leu Lys Val Gly Asp Arg Ile Leu Ser Leu Asn Gly 50 55 60

Lys Asp Val Arg Asn Ser Thr Glu Gln Ala Val Ile Asp Leu Ile Lys 65 70 75 80

Glu Ala Asp Phe Lys Ile Glu Leu Glu Ile Gln Thr Phe Asp Lys 85 90 95

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Ile Gln Gly Gln Tyr Pro Glu Ile Asp Ser Lys Leu Gln Arg Gly Asp 35 40 45

Ile Ile Thr Lys Phe Asn Gly Asp Ala Leu Glu Gly Leu Pro Phe Gln 50 60

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Val Thr Arg Pro Lys Pro

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Val Ala Gly Val Asp Pro Asn Gly Ala Leu Gly Ser Val Asp Ile Lys
Pro Gly Asp Glu Ile Val Glu Val Asn Gly Asn Val Leu Lys Asn Arg
Cys His Leu Asn Ala Ser Ala Val Phe Lys Ser Val Asp Gly Asp Lys
Leu Val Met Ile Thr Ser Arg Arg Lys
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Gly Cys Val Ile Thr His Val Tyr Pro Glu Gly Gln Val Ala Ala Asp

Lys Arg Leu Lys Ile Phe Asp His Ile Cys Asp Ile Asn Gly Thr Pro

Ile His Val Gly Ser Met Thr Thr Leu Lys Val His Gln Leu Phe His

Thr Thr Tyr Glu Lys Ala Val Thr Leu Thr Val Phe Arg Ala Asp Pro 70

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import locolization sequence targeting nucleus <223>

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<223>
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Phe Arg Asn Ile Leu Arg Leu Gln Ser Thr
           20
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<211> 4
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      import locolization sequence targeting endoplasmic reticulum
<223>
<400>
      15
Lys Asp Glu Leu
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<212> PRT
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      artificial
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      (1)..(4)
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Cys Cys Xaa Xaa
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